



Markers of neutrophil extracellular traps predict adverse outcome in community-acquired pneumonia: secondary analysis of a randomised controlled trial

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Neutrophil extracellular traps predict higher risk for clinical instability and 30-day mortality in pneumonia http://ow.ly/Gxbh30iqbom

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ABSTRACT Neutrophil extracellular traps (NETs) are a hallmark of the immune response in inflammatory diseases. However, the role of NETs in community-acquired pneumonia (CAP) is unknown. This study aims to characterise the impact of NETs on clinical outcomes in pneumonia.

This is a secondary analysis of a randomised controlled, multicentre trial. Patients with CAP were randomly assigned to either 50 mg prednisone or placebo for 7 days. The primary end-point was time to clinical stability; main secondary end-points were length of hospital stay and mortality.

In total, 310 patients were included in the analysis. Levels of cell-free nucleosomes as surrogate markers of NETosis were significantly increased at admission and declined over 7 days. NETs were significantly associated with reduced hazards of clinical stability and hospital discharge in multivariate adjusted analyses. Moreover, NETs were associated with a 3.8-fold increased adjusted odds ratio of 30-day mortality. Prednisone treatment modified circulatory NET levels and was associated with beneficial outcome.

CAP is accompanied by pronounced NET formation. Patients with elevated serum NET markers were at higher risk for clinical instability, prolonged length of hospital stay and 30-day all-cause mortality. NETs represent a novel marker for outcome and a possible target for adjunct treatments of pneumonia.

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Introduction

As a severe disease with high frequencies of morbidity and mortality, community-acquired pneumonia (CAP) ranks as one of the leading causes of hospital admission and is responsible for major socioeconomic costs [1]. Despite advances in antibiotic treatment and adjunct measures such as corticosteroids, hospitalised patients have persistently high mortality rates in the range of 5–15% [2]. Limited respiratory function and pulmonary, cardiovascular or metabolic comorbidities increase the likelihood of a severe course of CAP and associated complications [3].

The immediate immune response to the inoculation, proliferation and spread of pathogenic microbes in the lower airways is primarily borne by polymorphonuclear neutrophil granulocytes (neutrophils), the most abundant leukocytes in humans. Once primed, they migrate from the blood vessels through the bronchial and alveolar interstitium to engage the offending organisms within the airspaces [4]. They also target microbes invading the pulmonary parenchyma or crossing into the bloodstream. The weapons at their disposal are the generation of reactive oxygen species, phagocytosis, release of granular enzymes and, as recently discovered, the extrusion of chromosomal DNA in reticular arrays termed neutrophil extracellular traps (NETs) [5]. In addition, they release chemokines and cytokines that recruit further cells of the innate and specific arms of the immune system, firing up inflammation that can be harmful and cause pulmonary dysfunction.

NETs consist of chromatin complexes of cell-free DNA (cfDNA) and histones, with attached neutrophil granular proteins (e.g. neutrophil elastase, myeloperoxidase (MPO) and cathepsin G), which trap and kill infectious microbes [5]. The process of NET formation entails a novel form of cell death that has been termed NETosis [6].

NETs have been extensively investigated in the laboratory and in inflammatory conditions such as pre-eclampsia, coagulation disorders, atherosclerosis and autoimmune diseases (e.g. rheumatoid arthritis, lupus erythematosus) [7, 8]. However, clinical data on the role of NETs in infectious diseases are limited to case series in septicaemia. The characteristics of neutrophil activation to generate NETs in the initiation and resolution of pulmonary inflammation during the course of lower respiratory tract infections including CAP remain unknown.

Local and systemic effects of inflammation giving rise to measurable changes of mediators in the blood may be clinically useful for the guidance of decisions on therapy and for predicting the outcome of severe lower respiratory tract infections. Therefore, we aimed to explore the pattern and temporal dynamics of markers of NET formation and their associations with disease severity and relevant outcome measures of CAP. To investigate these questions, we chose a well-defined cohort of patients with CAP from a previous multicentre study, in which we recently showed that adjunct corticosteroids meaningfully shorten the time to clinical stability (TTCS) and length of hospital stay compared with placebo [9].

Materials and methods

Study subjects and design

This is a secondary analysis of an investigator-initiated, double-blind, randomised, placebo-controlled multicentre trial, in which adjunct treatment with prednisone in CAP was investigated. Details of the trial design and results have been published [9].

In brief, consecutive patients (≥18 years) presenting with CAP were enrolled at emergency departments or medical wards of seven tertiary care hospitals in Switzerland within 24 h of presentation.

The conduct of the trial adhered to the Declaration of Helsinki and Good Clinical Practice guidelines, and ethical committees of all participating hospitals approved the study before patient recruitment. The trial was registered at ClinicalTrials.gov (identifier NCT00973154).

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Conflict of interest: F. Ebrahimi reports receiving grants from Research Funds of the University of Basel during the conduct of the study. S. Giaglis reports receiving grants from Kantonsspital Aarau Research Council during the conduct of the study and has a patent (US 20160061824 A1) pending. S. Hahn has a patent (US 20160061824 A1) pending. C.A. Blum reports receiving grants from Research Funds Endocrinology Kantonsspital Aarau during the conduct of the study. M. Christ-Crain reports receiving grants from the Swiss National Science Foundation, the Gottfried and Julia Bangerter-Rhyner Foundation and the Nora van Meeuwen Häfliger Foundation, during the conduct of the study. P. Hasler reports receiving grants from the Kantonsspital Aarau Research Council during the conduct of the study and has a patent (US 20160061824 A1) pending.

Methods

Eligible patients were randomly assigned (1:1 ratio) to receive either 50 mg prednisone or placebo daily for 7 days. Patients, treating physicians, investigators and data assessors were masked to treatment allocation.

Informed consent was obtained within 24 h of admission to hospital. All patients were treated according to international CAP consensus guidelines [10]. Baseline data included medical history, relevant comorbidities, clinical variables relating to pneumonia and all variables required for the calculation of the Pneumonia Severity Index (PSI) [11]. Blood samples obtained over the course of 7 days after admission were used to investigate the temporal dynamics of systemic NET surrogate markers. Clinical study data were gathered up to 30 days after admission. Structured follow-up telephone interviews for secondary outcomes after discharge were performed on day 30.

Analysis

The primary end-point was TTCS, defined as time to clinical stabilisation of vital signs at two consecutive measurements \geqslant 12 h apart [9]. Secondary end-points included time to effective hospital discharge, all-cause mortality, duration of intravenous and overall antibiotic treatment, and CAP complications (including recurrence, acute respiratory distress syndrome, empyema, nosocomial infections until day 30, severe adverse events possibly related to CAP, intensive care unit (ICU) admission and readmission to hospital).

Analysis of surrogates of NETs

Blood samples (plasma and serum) from each patient were collected upon emergency department admission and on days 3, 5 and 7, and processed as previously described [9].

Cell-free nucleosomes as surrogate of NETosis were measured in serum samples of the total study population using Human Cell Death Detection ELISA PLUS (Roche Diagnostics, Basel, Switzerland). The concentrations of neutrophil elastase and MPO were measured in sera and plasma by sandwich ELISA, utilising the Elastase/ α 1-PI complex ELISA kit (Calbiochem/EMD, Gibbstown, NJ, USA) and the human MPO ELISA kit (Hycult Biotech, Plymouth Meeting, PA, USA), respectively. To specifically characterise the source of cell-free nucleosomes as NET-associated MPO-DNA complexes, MPO-specific capture and subsequent DNA-specific detection antibodies were used as previously described [12]. For detailed methodology, see supplementary material [13–20].

Statistical analysis

Unless stated otherwise, categorical variables are expressed as number (percentage) and continuous variables as median (interquartile range (IQR)) or mean±standard deviation. In order to assess an integrated value of NET surrogates over 7 days, area under the concentration curve (AUC) values were calculated. For AUC calculations missing NETs, values were imputed using multiple imputation.

For temporal changes in biomarker levels, a multilevel mixed-effects linear regression model was applied. Associations of comorbidities with baseline levels of cell-free nucleosomes were analysed by univariate and multivariate linear regression models. For the primary end-point, we used a multivariate Cox proportional hazards model based on a binary outcome of achieving or not achieving clinical stability. For all secondary end-points, we calculated unadjusted and adjusted estimates of the effect size and corresponding 95% confidence intervals using linear, logistic or Cox proportional hazards regression as appropriate. All multivariate models were adjusted for the same variables: treatment group, patient age, sex, body mass index, smoking status, PSI and comorbidities (congestive heart failure, chronic kidney disease, cerebrovascular disease, peripheral artery occlusive disease, chronic obstructive pulmonary disease (COPD), asthma), as well as clinical variables (days of symptoms before admission, systemic inflammatory response syndrome (SIRS) criteria, C-reactive protein and white blood cell count). Kaplan–Meier curves were used to illustrate TTCS based on NET quartiles (highest quartile versus lower three quartiles).

All statistical analyses were performed using Stata version 14.2 (StataCorp, College Station, TX, USA) and tests were done at a two-sided 5% significance level with two-sided 95% confidence intervals.

Results

Patient characteristics

Overall, 310 patients were randomly selected from the total study population of 785 patients for the current analysis. Baseline characteristics of treatment and placebo subgroups are presented in table 1. The median age was 75 years and 64.2% were male. The burden of pre-existing pulmonary disease was relatively low, with a positive smoking history in 23.2% and a history of COPD in 17.4%. At hospital admission, the median duration of symptoms was 4 days and approximately one-fifth (21.9%) of the patients had been pre-treated with an antibiotic. Overall, 78.5% of the patients had two or more SIRS

TABLE 1 Baseline characteristics and clinical variables of enrolled patients

	Total	Prednisone	Placebo	p-value
Subjects	310	151	159	
General characteristics				
Age years	75.0 (63.0-83.0)	77.0 (63.0-84.0)	73.0 (62.0-83.0)	0.19
Male	199 (64.2)	96 (63.6)	103 (64.8)	0.83
BMI kg·m ⁻²	25.9 (23.1–29.3)	25.1 (22.5–28.1)	26.4 (23.7–30.2)	0.009
Ever-smoker	72 (23.2)	36 (23.8)	36 [22.6]	0.80
Pack-years	10.0 (0.0-40.0)	10.0 (0.0-40.0)	10.0 (0.0-40.0)	0.80
Comorbidities				
Diabetes mellitus	68 (21.9)	31 (20.5)	37 (23.3)	0.56
Insulin treatment	25 (8.1)	12 (7.9)	13 (8.2)	0.94
COPD	54 (17.4)	32 (21.2)	22 (13.8)	0.088
Asthma	13 (4.2)	5 (3.3)	8 (5.0)	0.45
Heart failure	62 (20.0)	31 (20.5)	31 (19.5)	0.82
Hypertension	176 (56.8)	88 (58.3)	88 (55.3)	0.60
Cerebrovascular disease	28 (9.0)	13 (8.6)	15 (9.4)	0.80
Peripheral artery occlusive disease	20 (6.5)	8 (5.3)	12 (7.5)	0.42
Renal insufficiency	107 (34.5)	53 (35.1)	54 (34.0)	0.83
Neoplastic disease	19 (6.1)	9 (6.0)	10 (6.3)	0.90
Steroid pre-treatment	7 (2.2)	2 (1.4)	5 (3.1)	0.43
Antibiotic pre-treatment	68 (21.9)	31 (20.5)	37 (23.3)	0.56
Clinical variables	00 (2,	0. (20.0,	0. (20.0)	0.00
Days with symptoms n	4.0 (2.0-7.0)	4.0 (2.0-7.0)	4.0 (2.0-7.0)	0.50
Systolic blood pressure mmHq	121.0 (109.0–139.0)	122.0 (109.5–137.0)	120.0 (109.0–140.0)	0.93
Diastolic blood pressure mmHq	66.0 (58.0–74.0)	65.0 (57.5–74.0)	67.0 (58.0–73.0)	0.77
Pulse beats·min ⁻¹	85.0 (74.0–98.0)	85.0 (76.0–97.0)	85.0 (72.0–99.0)	0.86
Respiratory frequency breaths·min ⁻¹	20.0 (18.0–24.0)	20.0 (18.0–24.0)	20.0 (16.5–24.5)	0.56
Temperature °C (in-ear)	38.0 (37.4–38.5)	38.0 (37.4–38.5)	38.0 (37.4–38.4)	0.52
SIRS points	00.0 (07.1. 00.0,	00.0 (07.1. 00.0,	33.3 (37.1. 33.1,	0.02
0	6 (1.9)	4 (2.6)	2 (1.3)	0.86
1	61 (19.7)	27 (17.9)	34 (21.4)	0.00
2	114 (36.8)	56 (37.1)	58 (36.5)	
3	91 (29.4)	45 (29.8)	46 (28.9)	
4	38 (12.3)	19 (12.6)	19 (11.9)	
PSI score#	22 (,	(,	,	
1	32 (10.3)	14 (9.3)	18 (11.3)	0.55
II	47 (15.2)	25 (16.6)	22 (13.8)	0.50
iii	65 (21.0)	29 (19.2)	36 (22.6)	0.46
IV	118 (38.1)	60 (39.7)	58 (36.5)	0.55
V	48 (15.5)	23 (15.2)	25 (15.7)	0.90
PSI points	94.7 (36.7)	95.6 (37.9)	93.8 (35.6)	0.66
Laboratory values	, (66,	70.0 (07.77)	, 516 (5515)	0.00
Procalcitonin ng·dL ⁻¹	0.4 (0.2-2.4)	0.6 (0.2–2.5)	0.4 (0.2–2.0)	0.67
C-reactive protein mg·L ⁻¹	153.6 (80.2–237.5)	153.0 (80.5–237.5)	154.0 (73.7–240.0)	0.98
White blood cell count g·L ⁻¹	11.9 (8.7–15.9)	12.3 (8.8–16.1)	11.6 (8.7–15.5)	0.27
Neutrophil granulocytes g·L ⁻¹	9.9 (7.0–13.2)	10.3 (7.0–13.4)	9.6 (6.9–13.1)	0.26
Fasting glucose mmol·L ⁻¹	6.6 (5.8–7.5)	6.4 (5.6–7.5)	6.6 (5.9–7.6)	0.50
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Data are presented as n, median (interquartile range) or n [%], unless otherwise stated. BMI: body mass index; COPD: chronic obstructive pulmonary disease; SIRS: systemic inflammatory response syndrome; PSI: Pneumonia Severity Index. #: the PSI is a clinical prediction rule to calculate the probability of morbidity and mortality in patients with community-acquired pneumonia [11]; PSI risk class I corresponds to age \leqslant 50 years and no risk factors (\leqslant 50 points), risk class II to 51–70 points, risk class III to 71–90 points, risk class IV to 91–130 points and risk class V to >130 points.

criteria, and about half (53.6%) had severe pneumonia classified in the high-risk PSI classes IV and V, respectively.

Temporal dynamics of NETs

At baseline, serum cell-free nucleosomes as surrogates of NET formation were highly increased $(2.67\pm1.32$ absorbance units (AU)) compared with values in healthy controls reported as \sim 0.2 AU [13, 15, 17, 18]. They subsided gradually until day 7 (day 3: 1.48 \pm 1.03 AU; p<0.0005; day 5: 1.19 \pm 0.87 AU; p<0.0005;

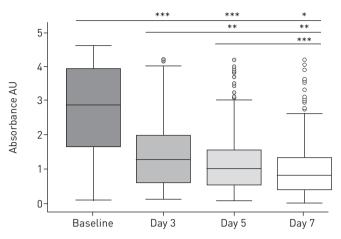


FIGURE 1 Box plots for levels of cell-free nucleosomes as a surrogate marker of neutrophil extracellular traps (NETs) in serum samples of the whole cohort (n=310) at four different time-points (baseline, and days 3, 5 and 7) in the course of community-acquired pneumonia. AU: absorbance units ($A_{405\,\mathrm{nm}}$ – $A_{490\,\mathrm{nm}}$). Each box signifies the upper and lower quartities, while the median is represented by a line within the box. Whiskers represent the upper and lower adjacent values; outliers are depicted as circles. At admission, mean levels of NETs were markedly increased (2.67±1.32 AU) and decreased gradually over time (day 3: 1.48±1.03 AU; p<0.0005; day 5: 1.19±0.87 AU; p<0.0005; day 7: 0.98±0.79 AU; p=0.042). Mixed-effects linear regression model. *: p<0.05; ***: p<0.005; ***: p<0.0005.

day 7: 0.98±0.79 AU; p=0.042; figure 1), when they were still markedly elevated compared with values of randomly selected healthy controls (characteristics of healthy controls are shown in supplementary table S1). In the plasma, cell-free nucleosomes were much lower and maximum values were reached at day 3 (data not shown). Quantification of MPO–DNA complexes showed a high correlation with values of cell-free nucleosomes, indicating that they represent NETs (r=0.55, R²=0.3, p=0.012) (supplementary figure S3).

Associations of initial NET surrogates with demographic characteristics, comorbidities and clinical variables

Multivariate linear regression models were used to investigate predictors of increased markers of NETosis at baseline. In univariate analysis, advanced age was associated with decreased levels of NETs (data not shown), although this association was no longer significant after full adjustment for multiple covariates (table 2). Smokers with a greater past exposure to tobacco, as measured by pack-years, had significantly lower NET levels on hospital admission ($-0.05~{\rm AU}~{\rm per}~10~{\rm pack}$ -years; p=0.042). There was likewise an inverse association of NET levels with heart failure. Patients with a history of heart failure had significantly lower NET levels on hospital admission compared with patients without heart failure ($2.33\pm1.45~{\it versus}~2.76\pm1.28~{\rm AU}$; p=0.036). This association remained robust in a fully adjusted model. As NETs are generated by neutrophils, NET levels were correspondingly associated with elevations of white blood cell counts as well as neutrophil counts (table 2).

Association between markers of NETs and disease outcome measures

Relevant disease outcome measures were analysed for NET surrogates at baseline as well as for the AUCs, integrating values of NET surrogates over 7 days. The primary end-point, *i.e.* TTCS, was significantly longer in patients from the highest AUC NET quartile (median (IQR) 5.0 (2.6-9.0) days) compared with the lower three quartiles (4.0 (2.0-7.4) days) with an adjusted hazard ratio (HR) of 0.97 (95% CI 0.94-0.99; p=0.041; figure 2 and table 3); a HR<1.0 corresponding to prolonged TTCS. For high baseline NET levels, a trend towards prolonged TTCS was noted (adjusted HR 0.91, 95% CI 0.82-1.01; p=0.088).

Median (IQR) time to effective discharge from hospital was longer in patients with the highest quartile of AUC NET levels (9.0 (5.0–14.0) days) compared with the three lower quartiles (7.0 (5.0–11.0) days) with an adjusted HR of 0.90 (95% CI 0.82–0.99; p=0.042). In fully adjusted logistic regression analyses, increased NET levels at admission were associated with a 3.8-fold odds ratio of 30-day mortality. In addition, a stepwise increase in mortality was found with mounting quartiles of NETs.

Higher markers of NETs at baseline were associated with a 0.52-day prolongation of i.v. antibiotic treatment (regression coefficient 0.52, 95% CI 0.1–0.94; p=0.015). There was no association of NETs with a composite end-point of complications connected with CAP or the probability of ICU admission (table 3).

TABLE 2 Association of baseline neutrophil extracellular trap (NET) levels with demographic characteristics, comorbidities and clinical variables

	Adjusted (multivariate) linear regression coefficient (95% CI)	p-value
General characteristics		
Age at study entry	-0.001 (-0.02-0.006)	0.292
Male	0.118 (-0.22-0.463)	0.502
BMI	-0.001 (-0.03-0.026)	0.892
Ever-smoker	0.147 (-0.23-0.529)	0.450
Pack-years (per 10 years)	-0.05 (-0.100.00)	0.042
Comorbidities		
Diabetes mellitus	0.160 (-0.23-0.559)	0.428
Insulin treatment	0.318 (-0.25-0.895)	0.279
COPD	-0.01 (-0.42-0.395)	0.943
Asthma	-0.24 (-0.99-0.504)	0.522
Heart failure	-0.46 (-0.890.02)	0.036
Hypertension	0.041 (-0.28-0.370)	0.802
Cerebrovascular disease	-0.23 (-0.79-0.326)	0.410
Peripheral artery occlusive disease	-0.26 (-0.93-0.409)	0.442
Renal insufficiency	-0.04 (-0.38-0.290)	0.775
Neoplastic disease	-0.36 (-1.0-0.283)	0.270
Clinical variables		
Steroid pre-treatment	0.327 (-0.41-1.07)	0.385
Antibiotic pre-treatment	-0.02 (-0.40-0.355)	0.891
Days with symptoms	0.004 (-0.01-0.021)	0.648
Temperature	-0.01 (-0.18-0.164)	0.902
Oxygen saturation	0.006 (-0.04-0.056)	0.790
PSI score	0.001 (-0.001-0.007)	0.886
Positive blood culture	0.115 (-0.38-0.615)	0.649
Pathogen detection	-0.05 (-0.41-0.312)	0.782
Neutrophil granulocytes	0.033 (0.006-0.061)	0.016
Leukocyte count	0.019 (-0.00-0.043)	0.116

Baseline NET levels were negatively associated with the quantity of pack-years and a history of heart failure, but positively associated with counts of neutrophils. p<0.05 statistically significant. BMI: body mass index; COPD: chronic obstructive pulmonary disease; PSI: Pneumonia Severity Index.

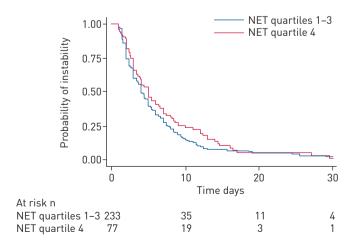


FIGURE 2 Kaplan–Meier curve of time to clinical stability (TTCS) for area under the concentration curves of neutrophil extracellular traps (NETs), an integrated value over 7 days: NET quartile 4 (highest quartile) *versus* NET quartiles 1–3. Higher NET levels are associated with prolonged TTCS (adjusted hazard ratio 0.97, 95% CI 0.94–0.99; p=0.041).

TABLE 3 Overview of primary and secondary end-points: neutrophil extracellular traps (NETs)

	Quartiles 1–3 AUC NETs#	Quartile 4 AUC NETs [¶]	NETs at baseline		AUC of NETs	
			Multivariate adjusted regression analysis: OR, HR, coefficient (95% CI)	p-value	Multivariate adjusted regression analysis: OR, HR, coefficient (95% CI)	p-value
Primary end-point						
TTCS days	4.0 (2.0-7.4)	5.0 (2.6-9.0)	0.91 (0.82-1.01)+	0.088	0.97 (0.94-0.99)+	0.041
Secondary end-points						
Time to effective hospital discharge days	7.0 (5.0–11.0)	9.0 (5.0–14.0)	0.90 (0.82-0.99)+	0.042	0.96 (0.94-0.99)+	0.012
Death (30 days)	9 (3.9)	4 (5.2)	3.81 (1.39-10.4) [§]	0.009	1.27 (1.05–1.54) [§]	0.012
Total duration of antibiotic treatment days	10.0 (7.0–12.0)	11.0 (9.0–14.0)	0.38 (-0.01-0.84) ^f	0.099	0.07 (-0.01-0.20) ^f	0.253
Intravenous antibiotic treatment days	5.0 (4.0-7.0)	6.0 (4.0–10.0)	0.52 (0.10–0.94) ^f	0.015	0.13 (0.01–0.24) ^f	0.021
CAP complications##	77 (33.1)	17 (22.1)	0.97 (0.78-1.22) [§]	0.853	0.99 (0.93-1.06)§	0.92
ICU admission	21 (9.0)	6 (7.8)	0.90 (0.58-1.39) [§]	0.645	0.97 (0.87–1.09) [§]	0.707

Data are presented as n, median (interquartile range) or n (%), unless otherwise stated, and adjusted for multiple variables (see Materials and methods). AUC: area under the concentration curve; TTCS: time to clinical stability; CAP: community-acquired pneumonia; ICU: intensive care unit. $^{\#}$: n=233; $^{\$}$: n=77; * : hazard ratio; $^{\$}$: odds ratio; $^{\$}$: regression coefficient; $^{\#}$ #: CAP complications: recurrence, acute respiratory distress syndrome, empyema, nosocomial infections until day 30, serious adverse events possibly related to CAP, ICU stay, readmission to hospital. p<0.05 statistically significant.

Effects of corticosteroids on NETs

In accordance with the main results from the original cohort, randomised treatment with prednisone over 7 days was associated with a significantly shorter median (IQR) TTCS (3.4 (2.0-6.0) *versus* 5.4 (3.0-9.0) days) and length of hospital stay (7.0 (5.0-11.0) *versus* 8.0 (5.0-13.0) days).

Corticosteroid treatment correlated with a slower decrease in levels of NET surrogates over time (figure 3). While there was no significant difference in mean values of cell-free nucleosomes as NET markers up to 3 days, corticosteroid treatment correlated with significantly higher NET formation on day 5 compared with placebo (1.09±0.88 AU in placebo group *versus* 1.31±0.84 AU in steroid group; p=0.023) and day 7 (0.79±0.74 *versus* 1.19±0.80 AU; p<0.0005). However, overall AUC values were not influenced by adjunct corticosteroid treatment compared with placebo (11.27±4.7 *versus* 10.74±5.1 AU; p=0.267). Likewise, patients from the placebo group presenting stable or rising markers of NETs after day 3 had higher hazards of clinical stability (adjusted HR 1.75, 95% CI 1.06–2.89; p=0.027).

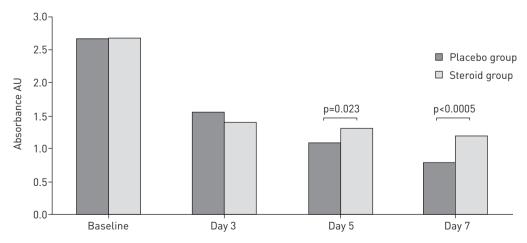


FIGURE 3 Mean absorbance of neutrophil extracellular traps (NETs): placebo group *versus* steroid group. AU: absorbance units ($A_{405 \text{ nm}}$ – $A_{490 \text{ nm}}$). On day 5 (1.09±0.88 *versus* 1.31±0.84 AU, respectively; p=0.023) and day 7 (0.79±0.74 *versus* 1.19±0.80 AU, respectively; p<0.0005) prednisone treatment was associated with significantly higher NET formation compared with placebo.

With regard to the beneficial effects of prednisone on clinical outcomes, we noted a significant effect modification by NET markers on TTCS ($p_{interaction}$ =0.008). Patients with levels of NETs in the highest quartile shared a 1.8-fold higher adjusted HR of clinical stability upon prednisone treatment compared with patients in the three lower quartiles (adjusted HR 1.79, 95% CI 1.03–3.12).

Discussion

The key findings of this study on NETs in patients with CAP are 3-fold. First, serum levels of cell-free nucleosomes as a surrogate marker for NETs are vastly increased in the context of CAP and remain at remarkably elevated levels even after 7 days. As the values for the ELISA of nucleosomes reflect a logarithmic scale, the actual magnitude of NET formation will be inclined to be underestimated, as it is in the range of mean elevations to the power of two to even four when compared with healthy volunteers [13, 15, 17, 18] and is considerably higher than that observed in autoimmune diseases such as rheumatoid arthritis with well-controlled activity [13]. While NETs are formed to fight infectious agents by entrapping and killing a wide variety of bacteria [21] and other microbes [22, 23], excessive activation of neutrophils and consecutive production of NETs are associated with the pathology of a number of airway disorders [24], such as COPD [25–27], impairment of alveolar gas exchange, lung dysfunction [28–31], and ultimately increased risk of early and late death.

Additionally, as measurements of NET surrogate markers were performed in serum samples, the results not only represent pre-formed systemic NETs, but also allow estimation of the propensity of pre-activated neutrophils to release NETs during coagulation. The magnitude of NETosis measured in the serum at baseline correlates with neutrophil and leukocyte cell count, but was reduced with the presence of heart failure. This observation may be due to pooling of neutrophils in the pulmonary circulation on account of reduced cardiac output [32]. While nicotine has been described to be a potent inducer of NETosis [33–35], in our cohort the number of pack-years was negatively associated with baseline levels of NETs, which could possibly be explained by exhaustion due to chronic activation of NETosis-related pathways.

The second important finding is that high levels of NETs at admission as well as a high integral of NETs during the course of CAP over 7 days (AUC) were strong independent risk factors for adverse outcome. Patients in the highest quartile had a median prolongation of TTCS by 1 day and of the length of hospital stay by 2 days, accompanied by increased duration of *i.v.* antibiotic treatment by 0.5 days. The differences in TTCS seemed to be less prominent in patients with very short (<5 days) or prolonged (>15 days) TTCS.

Remarkably, the results show that NET values at admission are associated with an almost 4-fold odds of all-cause mortality in CAP patients, representing a novel biomarker for outcome prediction, *i.e.* for identification of patients at risk.

These findings are consistent with previous observational studies that examined the association between cell-free DNA or nucleosomes and the risk of mortality in ICU patients. In a prospective observational cohort study from Finland on 225 patients with severe sepsis, admission concentrations of cfDNA were higher in ICU nonsurvivors compared with survivors [36]. Likewise, a retrospective observational study on 80 severe sepsis patients revealed high prognostic utility of cfDNA levels to predict ICU and hospital mortality [37]. Further observational studies support evidence of NETosis as a hallmark in the pathogenesis of systemic inflammation, hypercoagulability and sepsis [38–40]. These observations are in line with experimental data showing that intense pulmonary NET generation determines the disease severity in both viral and bacterial pneumonia [22, 41]. Seeking evidence that measured cell-free nucleosomes effectively mirrored NETs, we quantified complexes of MPO and DNA, measuring the typical components of NETs, DNA in combination with histones and neutrophil granule proteins. High correlations indicated that nucleosomes were, at the least, to a large extent derived from NET-forming neutrophils (supplementary figure S3).

Finally, our results suggest a bidirectional interaction of corticosteroids and NETs. While a 7-day treatment with prednisone was associated with a beneficial effect to achieve clinical stability, an overall reduction of length of hospital stay and a reduction in duration of *i.v.* antibiotic treatment [9], corticosteroids modulated NETosis, leading to a slower and more sustained decline of markers of NETs without increasing overall AUC after 3 days. Similarly, the subgroup of patients in the placebo group presenting with more sustained NETs levels from day 3 achieved clinical stability more rapidly than those with a faster decline. Accordingly, the temporal dynamics seem to exert a pivotal influence on clinical outcome: while excessive NETosis at baseline with consecutive exhaustion was associated with adverse outcome, moderate initial NET levels of sustained character were associated with a more beneficial outcome (supplementary figure S4). Against this background, beneficial effects of corticosteroids as adjunct therapy in CAP treatment may partly be driven by modulation of NET formation or pre-activation of neutrophils in the course of infection. In fact, significant effect

modification of corticosteroid effects on TTCS by NET levels supports this assumption. Yet, known effects of glucocorticoids on neutrophils prolonging the survival, increasing the mobilisation from the bone marrow and reducing marginalisation in the vasculature could in part account for enhanced NETosis [42, 43]. As there are no data on the effects of corticosteroids on NETosis to date, the underlying pathophysiological mechanisms are yet to be determined.

In conclusion, our findings suggest that cell-free nucleosomes serve as a surrogate marker of NETosis, which appears to be a highly dynamic and prognostic cellular antimicrobial defence mechanism in CAP. The data contribute fundamentally to the understanding of the pathophysiology of lower respiratory tract infections, which are largely driven by neutrophil activation. While NET formation is understandably necessary in microbial defence, modulators of NETosis may have potential therapeutic benefits to limit excessive neutrophil recruitment and activation while preserving host defence.

The main strengths of our study include: 1) the well-characterised cohort of patients with CAP of different severities representative for patients usually treated in emergency departments and hospitals, 2) the large number of patients, 3) the assessment of clinically relevant end-points, and 4) the high accuracy and reproducibility of the measurement methods. However, limitations that need to be considered include: 1) this was a secondary analysis of a randomly selected subgroup, 2) measured cell-free nucleosomes may not only mirror NET formation but could partly reflect other mechanisms of cell death such as necrosis or necroptosis, 3) quantified markers of NETs could be overestimated due to possible fragmentation and 4) the results may not be applicable to patients with less severe CAP, who do not require hospitalisation.

Further studies are required to characterise NETosis in lower respiratory tract infections and to determine the relative importance to the progression and remission of the condition, and whether targeting this pathway would be of therapeutic benefit to patients' outcomes.

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